

GenCore version 4.5  
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OM protein - protein search, using sw model

Run on: May 8, 2002 11:06:25 ; Search time 14.94 Seconds  
(without alignments)  
801.322 Million cell updates/sec

Title: US-09-911-513-2  
Perfect score: 2758  
Sequence: 1 MKRDHHHHQDKKTMNNEE.....MLGWHTRPLIATSMAKLSTN 532

Scoring table: BLUSUM62  
Gapop 10.0 , Gapext 0 .5

Searched: 212252 seqs, 22503292 residues

Total number of hits satisfying chosen parameters: 212252

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued\_Patents\_AA:  
 1: /cgns\_6/prodata/2/1aa/5A\_COMB.pep:  
 2: /cgns\_6/prodata/2/1aa/5B\_COMB..pep:  
 3: /cgns\_6/prodata/2/1aa/6A\_COMB..pep:  
 4: /cgns\_6/prodata/2/1aa/6B\_COMB..pep:  
 5: /cgns\_6/prodata/2/1aa/PCTMS\_COMB..pep:  
 6: /cgns\_6/prodata/2/1aa/backfile1..pep:  
 \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No. Score Query Match Length DB ID Description

Result No.	Score	Query Match	Length	DB ID	Description
1	2758	100.0	532	4 US-09-117-853-2	Sequence 2, Appli
2	1338.5	48.5	282	4 US-09-117-853-8	Sequence 8, Appli
3	1304.5	47.3	259	4 US-09-117-853-6	Sequence 6, Appli
4	1108.5	40.2	221	4 US-09-117-853-4	Sequence 4, Appli
5	713.5	25.9	166	4 US-09-117-853-10	Sequence 10, Appli
6	109.5	4.0	664	1 US-08-121-661-6	Sequence 6, Appli
7	104	3.8	2293	4 US-09-68-990-2	Sequence 2, Appli
8	101.5	3.7	924	3 US-08-619-812-8	Sequence 8, Appli
9	101.5	3.7	926	1 US-07-908-253-2	Sequence 2, Appli
10	101.5	3.7	926	1 US-08-455-970A-2	Sequence 2, Appli
11	101.5	3.7	926	1 US-08-387-156-6	Sequence 6, Appli
12	101.5	3.7	926	2 US-08-365-6	Sequence 6, Appli
13	101.5	3.7	926	2 US-08-878-748-6	Sequence 2, Appli
14	101.5	3.7	926	2 US-08-535-837-2	Sequence 6, Appli
15	101.5	3.7	926	3 US-09-124-491-6	Patent No. 5476657
16	101.5	3.7	926	6 5476657-3	Sequence 12, Appli
17	101.5	3.7	936	1 US-08-455-970A-12	Sequence 6, Appli
18	101.5	3.7	943	1 US-08-355-970A-10	Sequence 6, Appli
19	101.5	3.7	951	1 US-08-455-970A-14	Sequence 14, Appli
20	101.5	3.7	977	1 US-08-387-156-8	Sequence 8, Appli
21	101.5	3.7	977	2 US-08-694-365-8	Sequence 8, Appli
22	101.5	3.7	977	2 US-08-878-748-8	Sequence 8, Appli
23	101.5	3.7	977	3 US-09-124-491-8	Sequence 8, Appli
24	101.5	3.7	1069	1 US-07-177-315-9	Sequence 9, Appli
25	101.5	3.7	1069	1 US-08-170-126-4	Sequence 4, Appli
26	101.5	3.7	1069	3 US-08-954-118-4	Sequence 4, Appli
27	101.5	3.7	1098	1 US-07-777-15-7	Sequence 7, Appli

## ALIGNMENTS

RESULT 1  
US-09-117-853-2  
; Sequence 2, Application US/09117853  
; Patent No. 6307116  
; GENERAL INFORMATION:  
; APPLICANT: Harbord, Nicholas P  
; APPLICANT: Peng, Jinrong  
; APPLICANT: Carol, Pierre  
; APPLICANT: Richards, Donald E  
; TITLE OF INVENTION: Nucleic acid encoding GAI gene of Arabidopsis thaliana  
; FILE REFERENCE: 620-45  
; CURRENT APPLICATION NUMBER: US/09-117, 853  
; CURRENT FILING DATE: 1998-08-12  
; EARLIER APPLICATION NUMBER: PCT/GB97/00390  
; EARLIER FILING DATE: 1997-02-12  
; EARLIER APPLICATION NUMBER: GB 9602796. 6  
; NUMBER OF SEQ ID NOS: 12  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 2  
; LENGTH: 532  
; TYPE: PRT  
; ORGANISM: Arabidopsis thaliana  
; US-09-117-853-2

Query Match Best Local Similarity 100.0%; Score 2758; DB 4; Length 532;  
Mismatches 0; Indels 0; Gaps 0;

Qy 1 MKRDHHHHQDKKTMNNEE.....MLGWHTRPLIATSMAKLSTN 60  
Db 1 MKRDHHHHQDKKTMNNEE.....MLGWHTRPLIATSMAKLSTN 60

Qy 1 SNOGGGDYTTNKRKCNGVYETTATAESFRHVLVDQSENGVRVHLAACAVQ 180  
Db 1 SNOGGGDYTTNKRKCNGVYETTATAESFRHVLVDQSENGVRVHLAACAVQ 180

Qy 1 KENLTVAEALVKQIGFLAVSQIGAMRKVATYFAEALARRYRSPSOPIDHSLSDFLQW 240  
Db 1 KENLTVAEALVKQIGFLAVSQIGAMRKVATYFAEALARRYRSPSOPIDHSLSDFLQW 240

Qy 1 HFYETCPYLKFAHETANQAILAEFGQKKRVRVHVIDFMSMQLQMPALQA 300  
Db 1 HFYETCPYLKFAHETANQAILAEFGQKKRVRVHVIDFMSMQLQMPALQA 300

QY 301 RLTGIGPPADNFEDYLHEVGCKLAHLAEAHVEFEYRGFVANTLADDSMLERPSE 360  
 | ||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 301 RLTGIGPPADNFEDYLHEVGCKLAHLAEAHVEFEYRGFVANTLADDSMLERPSE 360  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 361 SVAVNSVFELHKLLGRPGADKVLGVVNQIKPEITVVEQESNHNSPIFLDRFTESLYV 420  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 361 SVAVNSVFELHKLLGRPGADKVLGVVNQIKPEITVVEQESNHNSPIFLDRFTESLYV 420  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 421 STLFDSLEGPSQGDQVMRQICHNVACDGPDRYERHETLSQWRNFGSAGFAAA 480  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 421 STLFDSLEGPSQGDQVMRQICHNVACDGPDRYERHETLSQWRNFGSAGFAAA 480  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 481 HIGSNFKQASMLIALFNGGEGYRVEESDGCLMIGWHTPLIASTAKLSTN 532  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 481 HIGSNFKQASMLIALFNGGEGYRVEESDGCLMIGWHTPLIASTAKLSTN 532  
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 ; RESULT 2  
 ; Sequence 8, Application US/09117-853-8  
 ; Patent No. 6307126  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Harberd, Nicholas P  
 ; APPLICANT: Peng, Jinrong  
 ; APPLICANT: Carol, Pierre  
 ; TITLE OF INVENTION: Nucleic acid encoding GAI gene of Arabidopsis thaliana  
 ; FILE REFERENCE: 620-45  
 ; CURRENT APPLICATION NUMBER: US/09/117,853  
 ; CURRENT FILING DATE: 1998-08-12  
 ; EARLIER APPLICATION NUMBER: PCT/GB97/00390  
 ; EARLIER FILING DATE: 1997-02-12  
 ; EARLIER APPLICATION NUMBER: GB 9602796.6  
 ; EARLIER FILING DATE: 1996-02-12  
 ; NUMBER OF SEQ ID NOS: 12  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 8  
 ; LENGTH: 282  
 ; TYPE: PRT  
 ; ORGANISM: Arabidopsis thaliana  
 ; US-09-117-853-8

Query Match 48.5%; Score 1338.5; DB 4; Length 282;  
 Best Local Similarity 94.0%; Pred. No. 1.e-120; Indels 17; Gaps 1;  
 Matches 264; Conservative 0; Mismatches 0; Indels 17; Gaps 1;  
 RESULT 4  
 US-09-117-853-4  
 ; Sequence 4, Application US/09117853  
 ; Patent No. 6307126  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Harberd, Nicholas P  
 ; APPLICANT: Peng, Jinrong  
 ; APPLICANT: Carol, Pierre  
 ; TITLE OF INVENTION: Nucleic acid encoding GAI gene of Arabidopsis thaliana  
 ; FILE REFERENCE: 620-45  
 ; CURRENT APPLICATION NUMBER: US/09/117,853  
 ; CURRENT FILING DATE: 1998-08-12  
 ; EARLIER APPLICATION NUMBER: PCT/GB97/00390  
 ; EARLIER FILING DATE: 1997-02-12  
 ; EARLIER APPLICATION NUMBER: GB 9602796.6  
 ; EARLIER FILING DATE: 1996-02-12  
 ; NUMBER OF SEQ ID NOS: 12  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO: 4  
 ; LENGTH: 221  
 ; TYPE: PRT  
 ; ORGANISM: Arabidopsis thaliana  
 ; US-09-117-853-4  
 Query Match 40.2%; Score 1108.5; DB 4; Length 221;  
 ; Sequence 6, Application US/09117853  
 ; Patent No. 6307126  
 ; GENERAL INFORMATION:

QY 1 MKRDHHHHHQDKTMMMNNEDDGNGMDELLAVLGKVRSSSEMADAQKLEEVMSNSVQ 60  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 1 MKRDHHHHHQDKTMMMNNEDDGNGMDELLAVLGKVRSSSEMADAQKLEEVMSNSVQ 60  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 61 EDDLSQLATEVHYNPAELYWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQPAIDSASS 120  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 61 EDDLSQLATEVHYNPAELYWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQPAIDSASS 120  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 66 EDDLSQLATEVHYNPAELYWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQPAIDSASS 103  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 66 EDDLSQLATEVHYNPAELYWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQPAIDSASS 103  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 121 SNOGGGDTYTNNRKLSQNGVYETTTATAESTRHVVLVDQENGVRVLHALCAEAVQ 180  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 121 SNOGGGDTYTNNRKLSQNGVYETTTATAESTRHVVLVDQENGVRVLHALCAEAVQ 180  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 181 KENITVAEALVKQIGFLAVSQIGAMRKVATYFAEALARRYRLSPSQSPIDHSLSDTLQM 240  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 181 KENITVAEALVKQIGFLAVSQIGAMRKVATYFAEALARRYRLSPSQSPIDHSLSDTLQM 240  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 184 SNOGGGDTYTNNRKLCNSNGVETTTATAESTRHVVLVDQENGVRVLHALCAEAVQ 180  
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 Db 184 SNOGGGDTYTNNRKLCNSNGVETTTATAESTRHVVLVDQENGVRVLHALCAEAVQ 180  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 241 HEYETCPYLKFAHETANGAILEAQGKRVHYD 274  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 241 HEYETCPYLKFAHETANGAILEAQGKRVHYD 274  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 QY 244 HEYETCPYLKFAHETANGAILEAQGKRVHYD 257  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 Db 244 HEYETCPYLKFAHETANGAILEAQGKRVHYD 257  
 | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |  
 ; RESULT 3  
 US-09-117-853-6  
 ; Sequence 6, Application US/09117853  
 ; Patent No. 6307126  
 ; GENERAL INFORMATION:

NUMBER OF SEQUENCES: 6  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: David J. O.  
 STREET: 2029 Century  
 CITY: Los Angeles  
 STATE: California  
 COUNTRY: U.S.  
 ZIP: 90067  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC  
 SOFTWARE: PatentIn Re  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: 0  
 FILING DATE:  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Oldenkamp, David  
 REGISTRATION NUMBER:  
 REFERENCE/DOCKET NUMBER:  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 310-788-5020  
 TELEFAX: 310-277-1297  
 INFORMATION FOR SEQ ID NO:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 664 amino acid  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 COMMENTS: US-08-421-661-6

RESULT		7		
Qy	472	FG 473	US-09-368-590-2	
Db	651	FG 652	; Sequence 2, Application US/09368590	
			; Patent No. 6187563	
			; GENERAL INFORMATION:	
			; APPLICANT: Solimena, Michele	
			; TITLE OF INVENTION: INTERACTING POLYPEPTIDES FOR	
			; TITLE OF INVENTION: AUTOANTIGENS OF AUTOIMMUNE DISEASES	
			; FILE REFERENCE: 101918-200 (OCR-41)	
			; CURRENT APPLICATION NUMBER: US/09/368 590	
			; CURRENT FILING DATE: 1998-08-04	
			; EARLIER APPLICATION NUMBER: 60/095, 657	
			; EARLIER FILING DATE: 1998-08-07	
			; NUMBER OF SEQ ID NOS: 8	
			; SOFTWARE: FastSEQ for Windows Version 3.0	
			; SEQ ID NO: 2	
			; LENGTH: 2293	
			; TYPE: PRT	
			; ORGANISM: Human	
			; US-09-368-590-2	
Query Match Score 3.8%; Best Local Similarity 19.9%; Pred. No. 2.1; Mismatches 93; Indels 130; Gaps 27;				
Matches 105; Conservative Matches 105; Gapless Matches 105;				
Qy	12	KKTMNMMNEEDGNG - MDELLAVLG - YKRSSEMADVAQKLEQLEVMMMSNVQE -----	61	27;
Db	1254	KKNQGLDRETOAHGPPLLEEVLERAGALASRSPEEAVRGLEQLOQASAWGLREAERQ	1313	
Qy	62	--DDLSQIATETVHYNPAELTYWLDMSL2DLNPFSSNAEYDLKAPGDAILNQFAIDSA	118	
Db	1314	QVLDAFOV - EQQYFDVAEAWL -----	11	
Qy	119	S --SSNQGGGGDTYTNNKLCKCSNGVVEPTTATAEST ---RHWVL -- VDSOENGV - 166	11	
Db	1350	KVRPSIGCGGGAGGGRCRRTYGA - CQDQEOSTLQLKKHQLEQGVENVEESIA	1407	
Qy	167	--RLVHALACAAEVQENILTVAAEVALKVQIGFLAVSQIGAMRKVA -- TYFAEALARRI	220	
Db	1408	QLSQRQRALLERGHP - DSEQ1SRQSQVDRI - YVALKELGERRVAYALEQQYW-----L	1458	
Qy	221	YRLSPSQSPDHSLSPTLQMHFYECPYL - KFAHFTANGAILEAQGKQRKVHVLFDSMS	278	
Db	1459	YQLSRQSELEHWIA -- EKEVVAQSPPELGODFEHVSVLQ -- ENFS -----EFASE	1504	
Qy	279	QGLQNPALMQALALRGPPVFRLTGIGPAPDNFDYLHVVGCKLAHLABAIIHVFIFYRG	338	
Db	1505	TGMAGERLARV-----	1	
		--NMQMDELICG ---HTARATMAWK ---		1538
Qy	339	FVANTLADJASMLLRPSETESVAVNSVFELKHKLGRPAIDKVLGVVNQIKPEIETVV	398	
Db	1539	--DGHNNEARVELLEMGTAAQSLASR - ELHKPFSDARELOGOTEKRRRLPRLTTP	1593	
Qy	399	EQSNNSNPFLDRFTESL HYYSTLFDSLLEGVPFGQDKTMSEVVLGKOCNVVACDGP	457	
Db	1594	EPRPSSA --MORTLRAFHDLQLQVSYORLQGAAQRLT - VYAGEHEAIAS -----	1644	
Qy	458	RVERHETLSQWRNRFSGAGAAHGSNA ---- FKQASMFLALFNG 4 99	1	
Db	1645	--REQEVLQGWKEILLSACEADARLHVSTADALRHSQVRDILLSWDG 1689	1	

QY 340 VANTIDLDLSM--LELPSEIBSVANSVFE----LHKLLRGAGIDKVLGVVNQI 390  
   |  |  |  |  |  |  |  |  
 Db 411 DARYLNQLDDNMKFLNLN-KELQARVIAITQQWDNNIGDLAGISRLGEKVL----- 463  
 Qy 391 KPEITVVQEESNHNSPIFLDRFTHYYSS--TLFDSLEGV---PSGQDKVMSEVYL 443  
 Db 464 -----SGKAYDAFFEGKHAKDLVQLDSANGITDVSNSGAKTQHILFR 509  
 Qy 444 GKQICQNVVACDGPDVERHET-----LSQWRNRFGSA-----GF 477  
 Db 510 -----TPLLPGTSHRERYQTGKEYITKLNLNIRDMSKITDGAASSTFDLTNVQRIGI 564  
 Qy 478 AAAHIGS-NAFKQASMLL-----FNGEGY-RVEESDG 510  
 Db 565 ELDNRGNVTKTKEKIIAKIGEQQDDNFVGSGCTEDGEGYDVRHYSRG 614

RESULT 9  
 US-07-908-253-2  
 ; Sequence 2, Application US/07908253  
 ; Patent No. 5534256  
 ; GENERAL INFORMATION:  
 ;   APPLICANT: POUTER, ANDREW A.  
 ;   APPLICANT: HARLAND, RICHARD J.  
 ; TITLE OF INVENTION: HAMOPHILUS SOMNUS OUTER MEMBRANE  
 ; TITLE OF INVENTION: PROTEIN EXTRACT ENRICHED WITH IRON-REGULATED PROTEIN  
 ; NUMBER OF SEQUENCES: 6  
 ; CORRESPONDENCE ADDRESS:  
 ;   ADDRESSEE: ROBERTA L. ROBINS  
 ;   STREET: 635 BRYANT STREET  
 ;   CITY: PALO ALTO  
 ;   STATE: CALIFORNIA  
 ;   COUNTRY: UNITED STATES OF AMERICA  
 ;   ZIP: 94301  
 ; COMPUTER READABLE FORM:  
 ; MEDIUM TYPE: FLOPPY DISK  
 ; COMPUTER: IBM PC compatible  
 ; OPERATING SYSTEM: PC-DOS/MS-DOS  
 ; SOFTWARE: Patentin Release #1.0, Version #1.25  
 ; CURRENT APPLICATION DATA:  
 ;   APPLICATION NUMBER: US/07/908,253  
 ;   FILING DATE: 19920702  
 ; CLASSIFICATION: 420  
 ; ATTORNEY/AGENT INFORMATION:  
 ;   NAME: ROBINS, ROBERTA L.  
 ;   REGISTRATION NUMBER: 33,208  
 ;   REFERENCE DOCKET NUMBER: 9000-0026  
 ; TELECOMMUNICATION INFORMATION:  
 ;   TELEPHONE: (415) 617-8999  
 ;   TELEFAX: (415) 327-3331  
 ; INFORMATION FOR SEQ ID NO: 2:  
 ; SEQUENCE CHARACTERISTICS:  
 ; LENGTH: 926 amino acids  
 ; TYPE: AMINO ACID  
 ; TOPOLOGY: Linear  
 ; MOLECULE TYPE: protein  
 ; US-07-908-253-2

Query Match 3.7%; Score 101.5; DB 1; Length 926;  
 Best Local Similarity 18.8%; Pred. No. 0.85; Mismatches 207; Indels 221; Gaps  
 Matches 122; Conservative 100; MisMatches 207; Indels 221; Gaps

Qy 19 EDDONGMDELLAV--LGYKVSESEMADVAQK-----LGQLEYMMSNQEDD 63  
 Db 28 DTEQGNGLQDVLKRAAEFLIEQPEREENIATQSLGTQIAIGTVAISLQVSAQIDK 87

Qy 64 LSQ-----ATEVHYHNPAAELYTWLDSMLTDLNPSSNAEYDKAIPGDAILNOFAI 115  
 Db 88 LLQKRTAGQALGSAESIVQNANAKTVLQSIQSLSGVLAGMDL-EALQNNs--NOHAL 144

Qy 116 DSASSNQGGGDTTYTWNKRLKCSGNVUETT-----A 144

145 AKAG------LELTNLSIENIANSVTKLDEGEQISQFGSKLNQTKGLGT 188  
 149 TAESTRHVYLVDSOENGVRVHLALLACAEVQKENITVAE--ALVKOIGF--LAWSQI 202  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 189 LGDKLKNICGLDKAGGLGDLVISLISGATAA--LVLADKNASTAKKVAGFELANQVV 244  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 203 GAMRK-VATYFAEALARRYLRLSPSOSPIDHSLSPDLMHFYETCPYL----KEAHFT 255  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 245 GNITKAVSSYY-LAQRTYAAGLSSTGPYALIASTVSL--AISPLAFAGIADRFNHAK 298  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 256 ANOALIEAF-----OGKRRVHYTDFMSMOSQLOPMLQALARPGG-----296  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 299 SLESYAAERFKKLGYGDNLIAEYORGTTDASVY-----AINPALAAJAGGVAAAAG 352  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 297 -----PPVFRLTGCGPPADNFYIHEVGCKLHLAEAH--YEEF-----YRGF 339  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 353 SVIASPIALLVSGITGVISTILOYSKO - AMFEHVNKTHNKIVWEKNNHGKNYFENGY 410  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 340 VANTLADDASM---LELRPSETESAVNSVF-----DHLKLURGPADKVLGVNNQI 390  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 411 DARYLANLQDNMKFLNN-KELQAOERVITAIIQQWDNNGLAGLISRGEKVL-----463  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 391 KPELFITVQEESNHSPIFLDRFTEFLHYSS--TLFDLSLEGV---PSGODKVMSEVYL 443  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 464 -----SGKAVDAFEGKHAKDKLVQLDSANGIIVVSNSGRAKTOHILFR 509  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 444 GKQICINVYACDGDPDRVERHET-----LSQMRNRFGSA-----GF 477  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 510 -----TPLTPGTSHREYRQTGKYEITKLININRDSWKTIDGAASSTFDTNVYQRIGI 564  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 478 AAAHIGS-NAFKQOASSMLAI-----FNGGGSY-RVEESDG 510  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 565 ELDNAGNYTKTKETKIIAKLGDDNVFVGSGTTEIDGGEGYDRVHYSRG 614  
 ; : : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :  
 RESULT 10  
 US-08-455-970A-2  
 Sequence 2, Application US/08455970A  
 Patent No. 5708155  
 GENERAL INFORMATION:  
 APPLICANT: POTTER, ANDREW A.  
 APPLICANT: REDMOND, MARK J.  
 APPLICANT: HUGHES, HOW P.A.  
 TITLE OF INVENTION: ENHANCED IMMUNOGENICITY USING LEUKOTOXIN  
 NUMBER OF SEQUENCES: 15  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: REED & ROBINS  
 STREET: 285 HAMILTON AVENUE, SUITE 200  
 CITY: PALO ALTO  
 STATE: CALIFORNIA  
 COUNTRY: UNITED STATES OF AMERICA  
 ZIP: 94301  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patentn Release #1.0, Version #1.25  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/455,970A  
 FILING DATE: 31-MAY-1995  
 CLASSIFICATION: 424  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 07/960,932  
 FILING DATE: 14-OCT-1992  
 ATTORNEY/AGENT INFORMATION:  
 NAME: ROBINS, ROBERTA L.  
 REGISTRATION NUMBER: 33,208  
 REFERENCE/DOCKET NUMBER: 9001-0016.10  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (415) 327-3400  
 TELEFAX: (415) 327-3231  
 INFORMATION FOR SEO ID NO:  
 1

SEQUENCE CHARACTERISTICS:  
 LENGTH: 926 amino acids  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-08-455-970A-2

Query Match Score 3.7%; Best Local Similarity 18.8%; Pred. No. 0.85; Matches 122; Conservative 100; Mismatches 207; Indels 221; Gaps 32;

Qy 19 EDDGNGMDLILAV--LGKVKRSSESMADYAKQ-----LEQLEVMMNSNVQEDD 63  
 Db 28 DTEQENGLQDLVKAABEGLTVEQRBERNNIATAQNSLGLTQTAIGLTERGVILSAPQIDK 87

Qy 64 LSQ-----ATETVHYNPAELYWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQFAI 115  
 Db 88 LLQKTKAGQALGSASSTIVONAKAKTVLSCIQTLSVLAGMDLD-EAQNNS-NQHAL 144

Qy 116 DSASSSNQGGGDYTNNKRLKCSNGVVETT-----A 148  
 Db 145 AKAG-----ELTNSLIENANSVKTLDGFEOISQFGSKLQNQKGLGT 188

Qy 149 TAESTRHHVVLVDSEQNGVHLALLACADEVAKENLTVAE---ALVKOIGF--LAVSQI 202  
 Db 189 LGDKLUNIGGIDKAGLGLDVSGLUSGATAA---LVLDKKNASTAKVKGAGFELANOVV 244

Qy 203 GAMRK-VATYFAEALARRYLSPSQSPIDHSLSDLTQMHFYETCPYL----KFAHFT 255  
 Db 245 GNTITAVSSYT---LAQRYAAGLSSGTGPVAALIASTVSL---AISPLAFAGIADKENHAK 298

Qy 256 ANQAILEAF-----QGKVKRHVIFDSNSQGLOWPALMOALRPGG-----296  
 Db 299 SLESYAERFKLKGYDGDNLLAJAYQRTGTIDASVT-----AINTALAAIAAGGVSAAAAG 352

Qy 297 -----PPVFRLTGIGPAPPDNFDLYHEVOCRAHLAIAETH---VFEP-----YRF 339  
 Db 353 SVIASPIALLYSGITGVISTILQYSQK- AMFEHVANKIHNKIVWEWNHGNKFENGY 410

Qy 340 VANTLADLDASM---LELRPSETESVAVNSFE-----LHKLIGRPGAIKDVKLVGVVNQI 390  
 Db 411 DARYLALNQDMKFLNLN-KELGAERTVATTOQWDDNNIGDLAGISRLGEKV-----463

Qy 391 KPEIPTVVEOSENHNINSPIFLDRETESLHYH---TLEDSLEGV---PSQDQKVMSEVYL 443  
 Db 464 -----SGKAVYDAFEEGKHAKDLVQDLSANGILDVNSGAKTQHILFR 509

Qy 444 GKQTCNVVACGDPVERVHET-----LSQWNRNFGSA-----GF 477  
 Db 510 -----TPLLTPGTBHRERVQTKGYEYITKLINIRVDSWKITDGASSTFDLTNVQORIGI 564

Qy 478 AAHIGS-NAFKQASMLL-----FNGGEGY-RVEEESDG 510  
 Db 565 ELDNAGNVTKTKETKIIAKLGEQDDNNVFVGSGTEIDGGEGYDRVHYSRG 614

RESULT 11  
 US-08-387-156-6  
 Sequence 6, Application US/08387156  
 Patent No. 5723129  
 GENERAL INFORMATION:  
 APPLICANT: POTTER, ANDREW A.  
 REDMOND, MARK J.  
 APPLICANT: HUGHES, HOW P. A.  
 TITLE OF INVENTION: Gonr-LEUKOTOXIN CHIMERAS  
 NUMBER OF SEQUENCES: 28  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: REED & ROBLINS  
 STREET: 635 BRYANT STREET  
 CITY: PALO ALTO  
 STATE: CALIFORNIA  
 COUNTRY: UNITED STATES OF AMERICA

ZIP: 94301  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: FLOPPY disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patentin Release #1.0, Version #1.25  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/387,156  
 FILING DATE: 10-FEB-1995  
 CLASSIFICATION: 424  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 07/960,932  
 FILING DATE: 14-OCT-1992  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 07/779,171  
 FILING DATE: 16-OCT-1991  
 ATTORNEY/AGENT INFORMATION:  
 NAME: ROBLINS, ROBERTA L.  
 REGISTRATION NUMBER: 33,208  
 REFERENCE/DOCKET NUMBER: 9001-0016.21  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (415) 617-8999  
 FAX: (415) 327-3231  
 INFORMATION FOR SEQ ID NO: 6:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 926 amino acids  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-08-387-156-6

Query Match Score 3.7%; Best Local Similarity 18.8%; Pred. No. 0.85; Matches 122; Conservative 100; Mismatches 207; Indels 221; Gaps 32;

Qy 19 BEDDNGMDLILAV--LGKVKRSSESMADYAKQ-----LGKVKRSSEMAVYAK-----LEQLEVMMNSNVQEDD 63  
 Db 28 DTEQENGLQDLVKAEEELGIEVQEERNNIATAQTSLGTQTAIGLTERGVILSAPQIDK 87

Qy 64 LSQ-----ATETVHYNPAELYTWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQFAI 115  
 Db 28 DTEQENGLQDLVKAEEELGIEVQEERNNIATAQTSLGTQTAIGLTERGVILSAPQIDK 87

Qy 64 LSQ-----ATETVHYNPAELYTWLDSMLTDLNPPSSNAEYDLKAIPGDAILNQFAI 115  
 Db 88 LLQKTKAGQALGSASSTIVONAKAKTVLSCIQTLSVLAGMDLD-EAQNNS-NQHAL 144

Qy 116 DSASSSNQGGGDYTNNKRLKCSNGVVETT-----A 148  
 Db 145 AKAG-----ELTNSLIENANSVKTLDGFEOISQFGSKLQNQKGLGT 188

Qy 149 TAESTRHHVVLVDSEQNGVHLALLACADEVAKENLTVAE---ALVKOIGF--LAVSQI 202  
 Db 189 LGDKLUNIGGIDKAGLGLDVSGLUSGATAA---LVLDKKNASTAKVKGAGFELANOVV 244

Qy 203 GAMRK-VATYFAEALARRYLSPSQSPIDHSLSDLTQMHFYETCPYL----KFAHFT 255  
 Db 245 GNTITAVSSYT---LAQRYAAGLSSGTGPVAALIASTVSL---AISPLAFAGIADKENHAK 298

Qy 256 ANQAILEAF-----QGKVKRHVIFDSNSQGLOWPALMOALRPGG-----296  
 Db 299 SLESYAERFKLKGYDGDNLLAJAYQRTGTIDASVT-----AINTALAAIAAGGVSAAAAG 352

Qy 297 -----PPVFRLTGIGPAPPDNFDLYHEVOCRAHLAIAETH---VFEP-----YRF 339  
 Db 353 SVIASPIALLYSGITGVISTILQYSQK- AMFEHVANKIHNKIVWEWNHGNKFENGY 410

Qy 340 VANTLADLDASM---LELRPSETESVAVNSFE-----LHKLIGRPGAIKDVKLVGVVNQI 390  
 Db 411 DARYLALNQDMKFLNLN-KELGAERTVATTOQWDDNNIGDLAGISRLGEKV-----463

Qy 391 KPEIPTVVEOSENHNINSPIFLDRETESLHYH---TLEDSLEGV---PSQDQKVMSEVYL 443  
 Db 464 -----SGKAVYDAFEEGKHAKDLVQDLSANGILDVNSGAKTQHILFR 509

Qy 444 GKQTCNVVACGDPVERVHET-----LSQWNRNFGSA-----GF 477  
 Db 510 -----TPLLTPGTBHRERVQTKGYEYITKLINIRVDSWKITDGASSTFDLTNVQORIGI 564

Qy 478 AAHIGS-NAFKQASMLL-----FNGGEGY-RVEEESDG 510  
 Db 565 ELDNAGNVTKTKETKIIAKLGEQDDNNVFVGSGTEIDGGEGYDRVHYSRG 614

RESULT 11  
 US-08-387-156-6  
 Sequence 6, Application US/08387156  
 Patent No. 5723129  
 GENERAL INFORMATION:  
 APPLICANT: POTTER, ANDREW A.  
 REDMOND, MARK J.  
 APPLICANT: HUGHES, HOW P. A.  
 TITLE OF INVENTION: Gonr-LEUKOTOXIN CHIMERAS  
 NUMBER OF SEQUENCES: 28  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: REED & ROBLINS  
 STREET: 635 BRYANT STREET  
 CITY: PALO ALTO  
 STATE: CALIFORNIA  
 COUNTRY: UNITED STATES OF AMERICA

RESULT 12  
us-08-694-865-6  
; Sequence 6, Application US/08694865  
; Patent No. 5837268  
; GENERAL INFORMATION:  
; APPLICANT: POTTER, ANDREW A.  
; APPLICANT: MANN, JOHN G.  
; TITLE OF INVENTION: GNRH-LEUKOTOXIN CHIMERAS  
; NUMBER OF SEQUENCES: 34  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: REED & ROBINS LLP  
; STREET: 285 HAMILTON AVENUE, SUITE 200  
; CITY: PALO ALTO  
; STATE: CA  
; ZIP: 94301  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/694,865  
; FILING DATE: 09-AUG-1996  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: MCCRAYCEN, THOMAS P.  
; REGISTRATION NUMBER: 38,548  
; REFERENCE/DOCKET NUMBER: 9001-0016.22  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (415)327-3400  
; TELEFAX: (415)327-3231  
; INFORMATION FOR SEQ ID NO: 6:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 926 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: Protein  
; US-08-694-865-6

Query Match 3.78; Score 101.5; DB 2; Length 926;  
Best Local Similarity 18.8%; pred. No. 0.85; Indels 221; Gaps 32;

Matches 122; Conservative 100; Mismatches 207; Indels 221; Gaps 32;

Qy 19 EDDGNGMDELLAV --LGKVKRSSEMDAQK-----LEQLEMMMSNVQEDD 63  
Db 28 DTEQNGLQDVKAAEELGVQEERERNTAATQTSLGTTQATGERTIVSAPQDK 87  
Qy 64 LSQ-----ATEVTHYPAELTYWLDQMLTDINPPSSNAEYDILKAFGDLNQFAI 115  
Db 88 LLQKTKAGQALGSALSIVQANKAKTVLSQSLIGSVLAGMDLD-EALQNN--NQHAL 144  
Qy 116 DSASSSNQGGGDTYTNNKRLKCSNGNVERTT-----A 148  
Db 145 AKAG -----LELTSLTIANSTYKTLDFGEQLSQFGKLQNFKGLP 188  
Qy 149 TAESTRHVVLYDSQENGVRVYHALLACAENVKENITVAE--ALVKQIGF--LAWSQI 202  
Db 189 LGDKLNKNGDKAQGLDGLDTSGLSGATA---LVLARKNASTAKVYTAGFELANQVY 244  
Qy 203 GAMRK-VATVFAEALARRYRPLSPQSPIHSLSDTLQMMPFYETCPYL-----KFAHFT 255  
Db 245 GNITRAVSSYI--LAQRVAGLSSTSHPVYALIASTVSL--AISPLAFAGIADKFNHAK 298

Qy 256 ANQAILEAF-----OGKKRVHVIDSMSOGLQWPALMOALALRPGG----- 296  
Db 299 SLESTYPERFKLKGYDGNLILAEYQRTGTIDASVT-----AINTLAATAGGYSAAAAG 352  
Qy 297 ----PPVFRLETGIGPAPDNFDLYHEVGCKLHLAEAH--VVEE-----YRGF 339  
Db 353 SVIASPIALLYSGTICVISTLQYSRQ--AMFEHVNKINHKIVEWKNNHGKNYFENGY 410  
Qy 340 VANTLADDASM---LELRSETESVAVNSVF----LHKLIGRPGADKVLGTVVNQI 390  
Db 411 DARYLFLNQDMKFLINLN-KELQARVIAITQQWDNNIGDLAGTSRGEKVL----- 463  
Qy 391 KPEIIFTVVEQESNHNSPIELDRFTESHYHS--TFLDSLEGV----PSGQDKVMSSEVYL 443  
Db 464 -----SGKAYDAFEEBKHKADKLVQDSSRAKTOHILFR 509  
Qy 444 GKQICMVVACDGPDVMVERHET-----LSQWRNRFGSA-----GF 477  
Db 510 -----TPLLTPGTETHERVQTGKYEVITKLNINRVDWKLTDGAASTFDLTNVYQRGIGI 564  
Qy 478 AAAHIGS-NAFKQASHMLAL-----FNGGEGY-RVEESDG 510  
Db 565 ELDNAGNVTKTKEKTIIAKLGEGDDNNFVGSGTTEIDGEGSYDRHYHSGR 614

RESULT 13  
US-08-878-748-6  
; Sequence 6, Application US/08878748  
; Patent No. 5869126  
; GENERAL INFORMATION:  
; APPLICANT: POTTER, ANDREW A.  
; APPLICANT: REDMOND, MARK J.  
; APPLICANT: HUGHES, HOW P.A.  
; TITLE OF INVENTION: Gnrh-leukotoxin chimeras  
; NUMBER OF SEQUENCES: 28  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: REED & ROBINS  
; STREET: 635 BRYANT STREET  
; CITY: PALO ALTO  
; STATE: CALIFORNIA  
; ZIP: 94301  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/878,748  
; FILING DATE: 19-JUN-1997  
; CLASSIFICATION: 536  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 08/387,156  
; FILING DATE: 10-FEB-1995  
; APPLICATION NUMBER: US 07/960,932  
; FILING DATE: 14-OCT-1992  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/779,171  
; FILING DATE: 16-OCT-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: ROBINS, ROBERTA L.  
; REGISTRATION NUMBER: 33,208  
; REFERENCE/DOCKET NUMBER: 9001-0016.21  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (415) 617-8999  
; TELEFAX: (415) 327-3231  
; INFORMATION FOR SEQ ID NO: 6:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 926 amino acids  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 926 amino acids  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; US-08-878-748-6

CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/535,837  
 FILING DATE: 27-SEP-1995  
 CLASSIFICATION: 424  
 ATTORNEY/AGENT INFORMATION:  
 NAME: ROBINS, ROBERTA L.  
 REGISTRATION NUMBER: 33,208  
 REFERENCE/DOCKET NUMBER: 9001-0026.01  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (415) 327-3400  
 TELEFAX: (415) 327-3231  
 INFORMATION FOR SEQ ID NO: 2:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 926 amino acids  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-08-535-837-2

---

Query Match Score 3.7%; Length 926;  
 Best Local Similarity 18.8%; Pred. No. 0.85;  
 Matches 122; Conservative 100; Mismatches 207; Indels 221; Gaps 32;

Qy 19 FEDDGNGMDELLAV---LGYKVRSSSEMDAYAQK-----LEQLEVMMMSNVQEDD 63  
 Db 28 DTEQGNGLQDLVKAEEELGIVRERBNNTATAQTSLSGLTIQTAIGTERULVSLAPQIDK 87

Qy 64 LSQ-----ATEVHYNPAELYWLDSMLTDNPPSSNAEYDLKPGDAILNQFAI 115  
 Db 88 LLQKTKAGQALGSAESTVONRANKAKTVLSSLQSTILGSVLAGMDLD-EALQNS--NQHAL 144

Qy 116 DSASSSNQGGGDTYTNNKRKCSNSGVETT-----A 148  
 Db 145 AKAG-----LELTSLIENIANSVKTLDFFGEQISQFGSKLQNIKGLGT 188

Qy 149 TAESTRHVVLVDQENGVLRVHALLACAEEAVQKENLTVAE---ALVKQIGF---LAVSQI 202  
 Db 189 LGDKLNAGGIDKAGLGLDVTSGLUSGATA---LVLAQDNASTAKVKGAFELANQVV 244

Qy 203 GAMRK-VATYFAEALARRIYLSPSSOPIDHSLSPTDLOMHFYETCPYL----KFAHFT 255  
 Db 245 GNIITKAVSSYT---LAQRVAAGLSSGTGPVAALIASTVSL--AISPLAPAGIADKFENHAK 298

Qy 256 ANQAILEAF-----QGKRVHVLDIFSMSSQGLQWPALQALARPQG-----296

Db 299 SLESYAERFKKLGDGDNLLAEYQRTGTTDASVT-----AINTALAAIAAGGVSAAAAG 352  
 Qy 297 -----PPVFRLTGIGPAPNPNDYLYHEVGCKLHLAEATH---VFEFE-----YRGF 339  
 Db 353 SVIASPIALLSQTGIVSTLQLYSKQ---AMFEHVANKIHKNKIVEWEKNNHGKNWFENGY 410

Qy 340 VANTLADDASM---LELRPSEIESYAVNSYFE-----LHKLIGRPGLDVKLVGVNQI 390  
 Db 411 DARYLANLQDMKFLNLN-KELQERVRTATQQWDNNIGDLAGISRLGEKVL-----463

Qy 391 KPEIITVVEQFSNHNSPIFLDRFTESLHYS---TLFDSLGEV---PSGDQKYMSEVYL 443

Db 464 -----SGKAYVDAFEEGKHTRAKDLVYQOLDSANGIIIDVNSNSKAKTQHILPR 509  
 Qy 444 GKQICNVACDGPDRVERHET-----LSQWRNRFGSA-----GF 477

Db 510 -----TPLLTPGTEHRERVQTKYXYITKLINRNRYDSWKITDGAASSTFDTNVYQORIGI 564

Qy 478 AAHIGS-NAFKOASMLLA-----FNGGEGY-RVEESDG 510  
 Db 565 ELDNAGNVTKTKEKTIKAKLEGDDNNVFVGSGETIDGEGYDRVHYSRG 614

RESULT 14  
 US-08-535-837-2  
 Sequence 2 Application US/08535837  
 Application No. 5385289  
 GENERAL INFORMATION:  
 APPLICANT: POTTIER, ANDREW A.  
 APPLICANT: HARLAND, RICHARD J.  
 TITLE OF INVENTION: HAEMOPHILUS SOMNUS OUTER MEMBRANE  
 TITLE OF INVENTION: PROTEIN EXTRACT ENRICHED WITH IRON-REGULATED PROTEINS  
 NUMBER OF SEQUENCES: 6  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: ROBERTA L. ROBINS  
 STREET: 285 HAMILTON AVENUE, SUITE 200  
 CITY: PALO ALTO  
 STATE: CALIFORNIA  
 COUNTRY: UNITED STATES OF AMERICA  
 ZIP: 94301  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patentin Release #1.0, Version #1.25

RESULT 15  
 US-09-124-491-6  
 Sequence 6 Application US/09124491  
 Application No. 6022960

GENERAL INFORMATION:  
 APPLICANT: POTTER, ANDREW A.  
 APPLICANT: MANNES, JOHN G.  
 TITLE OF INVENTION: GnRH-LEUKOTOXIN CHIMERA  
 NUMBER OF SEQUENCES: 34  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: REED & ROBINS LLP  
 STREET: 285 HAMILTON AVENUE, SUITE 200  
 CITY: PALO ALTO  
 STATE: CA  
 COUNTRY: USA  
 ZIP: 94301  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: PatentIn Release #1.0, Version #1.30  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/09/124,491  
 FILING DATE:  
 CLASSIFICATION:  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 08/694,865  
 FILING DATE: 09-AUG-1996  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 08/387,156  
 FILING DATE: 10-FEB-1995  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 07/960,932  
 FILING DATE: 14-OCT-1992  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: US 08/779,171  
 FILING DATE: 16-OCT-1991  
 ATTORNEY/AGENT INFORMATION:  
 NAME: MCCRAYCEN, THOMAS P.  
 REGISTRATION NUMBER: 38,548  
 REFERENCE/DOCKET NUMBER: 9001-0016.22  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (415)327-3400  
 TELEFAX: (415)327-3231  
 INFORMATION FOR SEQ ID NO: 6:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 926 amino acids  
 TYPE: amino acid  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-09-124-491-6

Search completed: May 8, 2002, 11:08:17  
 Job time: 112 sec

Query Match Score 101.5; DB 3; Length 926;

Best Local Similarity 18.8%; Pred. No. 0.85; Matches 207; Indels 221; Gaps 32;

Matches 122; Conservative 100; Missmatches 207; Indels 221; Gaps 32;

Qy 19 EDDGNGMDELLAV--LGKVKRSEMDAYAQK-----LEQLEMMMSNVQEDD 63

Db 28 DTEQQNGLQDLVKAABELGLIEVQREERNNATAQTSLGTQTAIGLTERTGIVLSAQPQDK 87

Qy 64 LSQL-----ATEVHYNPAELTWDSLMLTDLNPPSSNAEYDLKAIGDAILNQFAI 115

Db 88 LLQKTKAGQALGSAESTVQVANKAKTVLQSIQLGSVLAQMDLD-EALQNN -NOHAL 144

Qy 116 DSASSSNQGGGDTYTNNKRLKCSNGVETT-----A 148

Db 145 AKAG-----DELTNSLIENIANSVKTLDERGEQISQFGSKLQNITKLGLFT 188

Qy 149 TAESTRHVVLYDQENGVRVLYHALLACAEAVQKENITVAE--ALVKGIF--LAVSQ1 202

Db 189 LGDKLKNIGGLDAGLGLDTISGLSGATAA---LYLAKDNNASTAKVAGFELANQVY 244

Qy 203 GAMRK-VATYFAEALARRYLRLSPSOSPIBHSLSLQMFYETCPY-----KFAHFT 255

Db 245 GNITRAVSSIT--LAQRAVAGLSSTGPVVALIASTVSL---AISPLAFFGIADEKFHNHK 298

Qy 256 ANQALEAF-----QGKRRVHVYDFSMSQGLQMPALMOALAALRPGG----- 296

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